



Supplemental Figure WWW1. A 13-exon Gene, *pfcrt*, is Located Within the 36 kb Segment Linked to CQR

The coding regions of the individual exons, arranged 5' to 3', have lengths of 91, 269, 173, 133, 72, 76, 83, 51, 57, 93, 45, 55 and 74 nucleotides. Crossover points (vertical arrows) delineate the boundaries of the CQR determinant previously mapped in the HB3×Dd2 genetic cross (Su *et al.*, 1997). Relative sizes and directions of the open reading frames are shown beneath the chromosome line.

Introns confirmed by RT-PCR are indicated for *cg7*, *hsp86* and *pfcrt*. C, ClaI; K, KpnI; L, Sall; M, BamHI; N, NcoI; V, PvuII.

Nucleotide distances along the chromosome line correspond to nucleotide positions of GenBank submission AF030694.